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Fig. 1A
18L



Fig. 1B
R18L

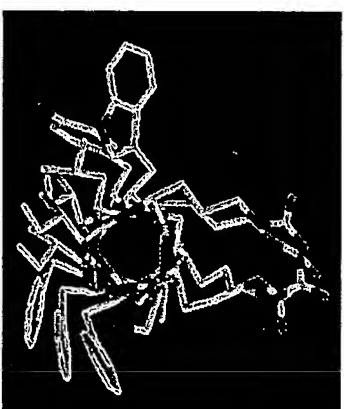


Fig. 1C
18A

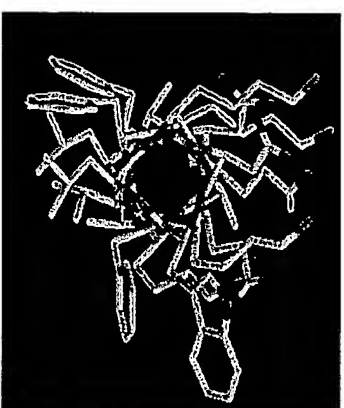


Fig. 1D

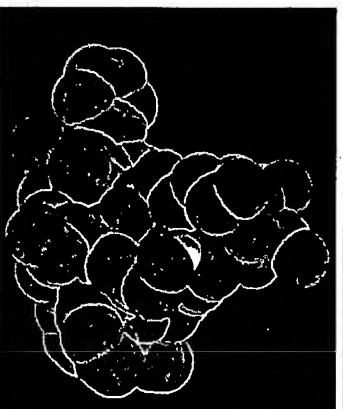


Fig. 1E

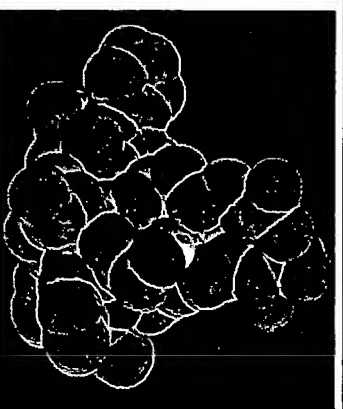


Fig. 1F



Figs. 1A-1F

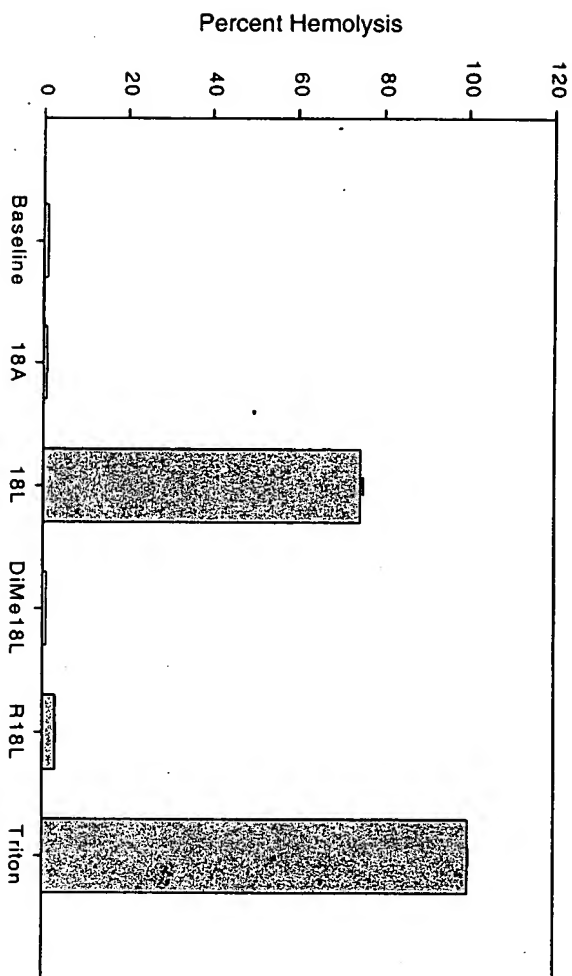
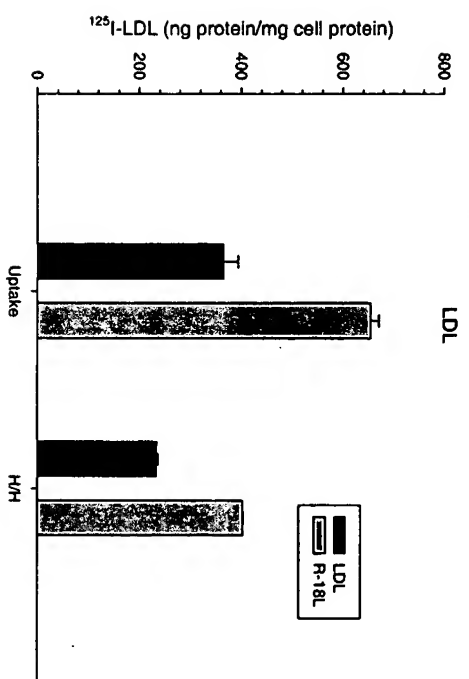
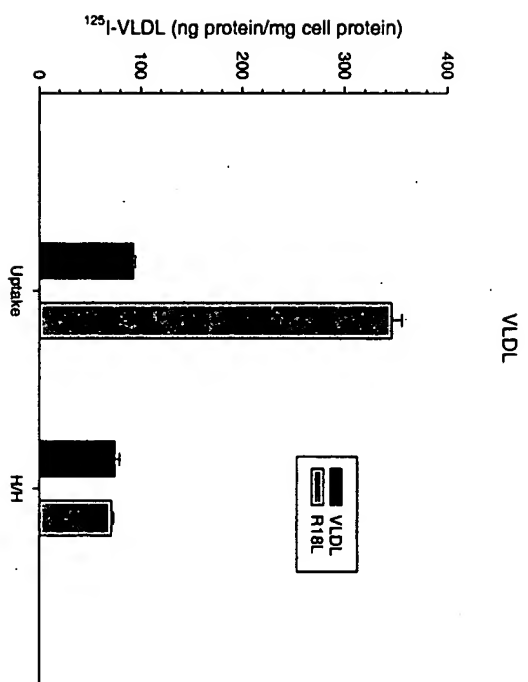


Fig. 2



Figs. 3A - 3B

Ac-R18L-NH₂ i.v. (100 µg/mouse)
Apo E null, fasted

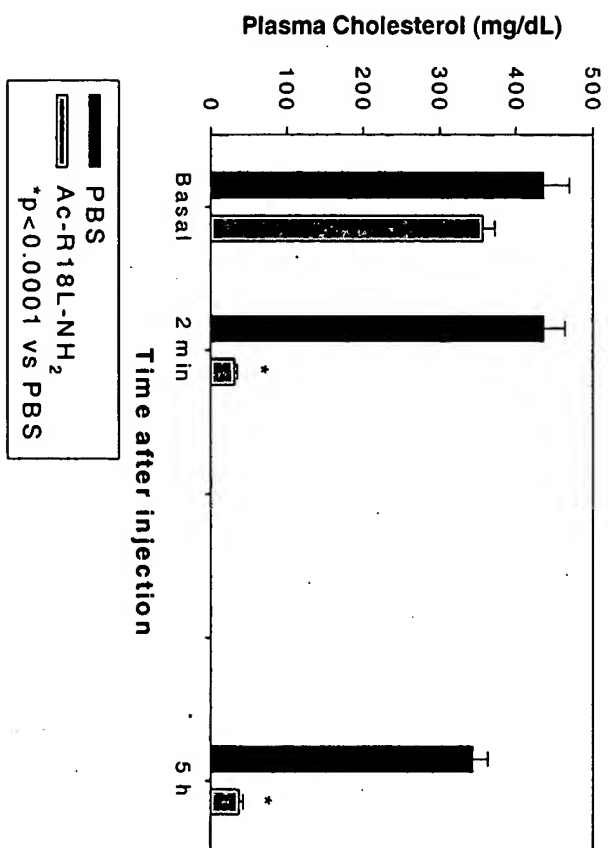


Fig. 4

Ac-R18L-NH₂, i.v. dose dependency
Apo E null; fasted

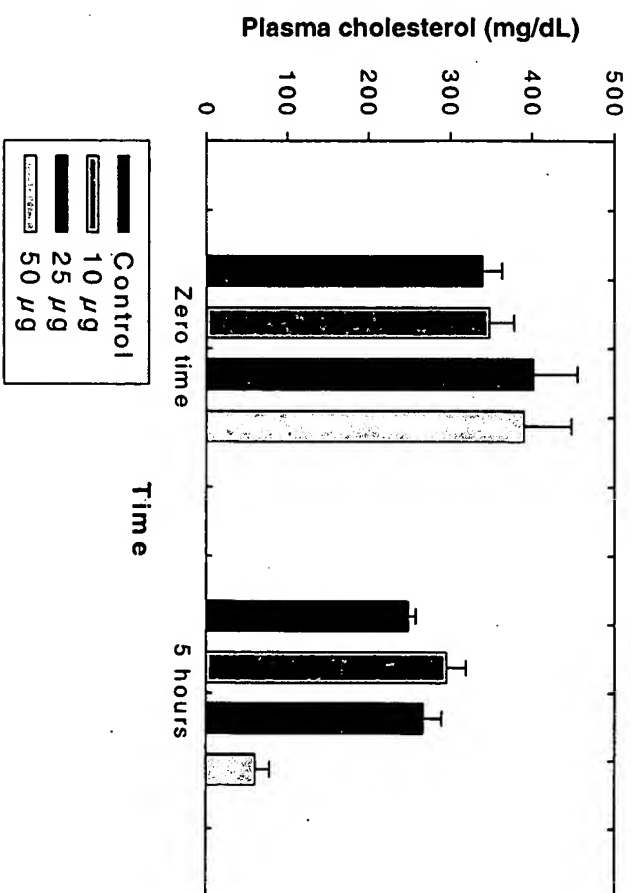


Fig. 5

24 h fecal total cholesterol
(collected from time of injection;
apo E null; 100 μ g R18L i.v.)

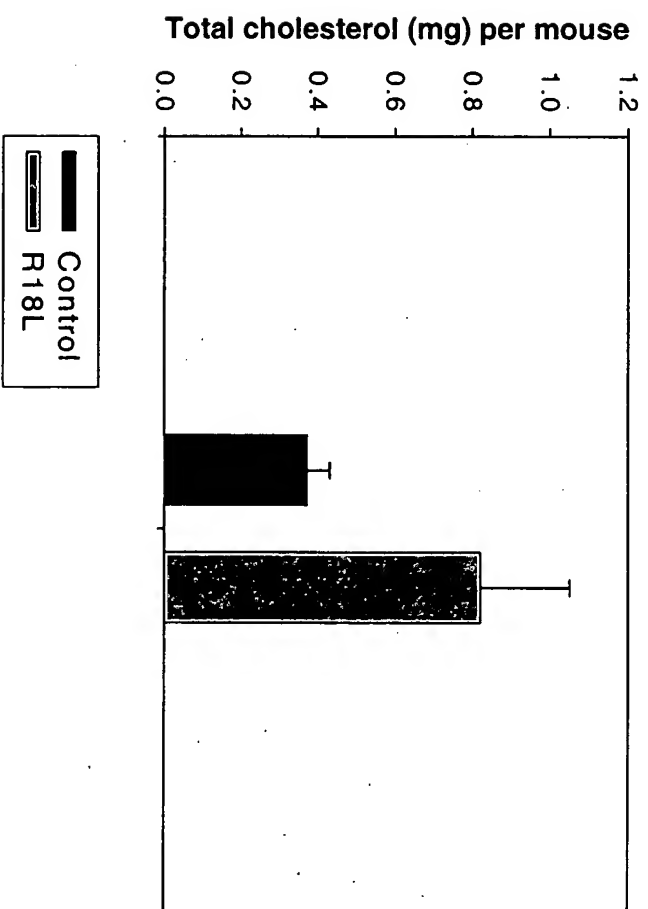


Fig. 6

In vitro mixing experiment
[125I]Ac-(R)18L-NH₂
LDLR^{-/-}, normal chow, fasted

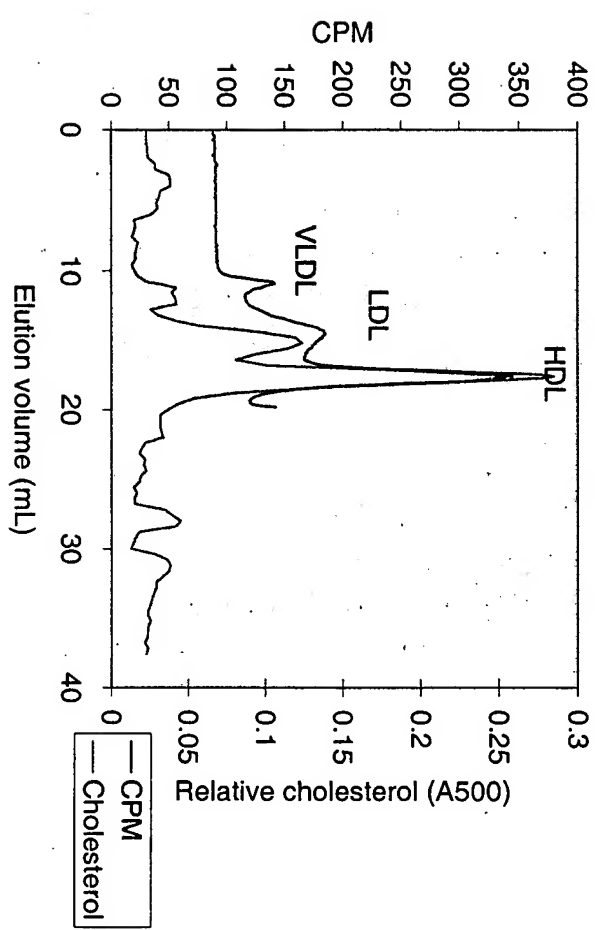
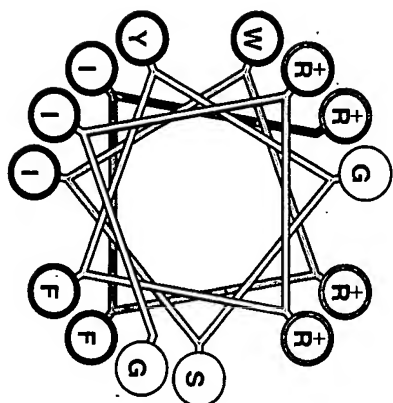
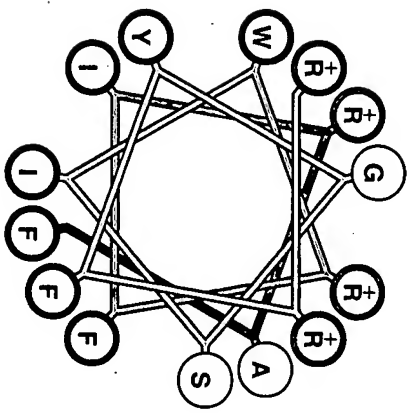


Fig. 7

1. A method of determining the relative
 2. positions of the various components of a
 3. system, comprising:
 4. (a) determining the relative positions of
 5. the various components of the system;
 6. (b) determining the relative positions of
 7. the various components of the system;
 8. (c) determining the relative positions of
 9. the various components of the system.



R14L-1
Fig. 8A



R14L-2
Fig. 8B

Uptake of LDL: Effect of R14L peptides

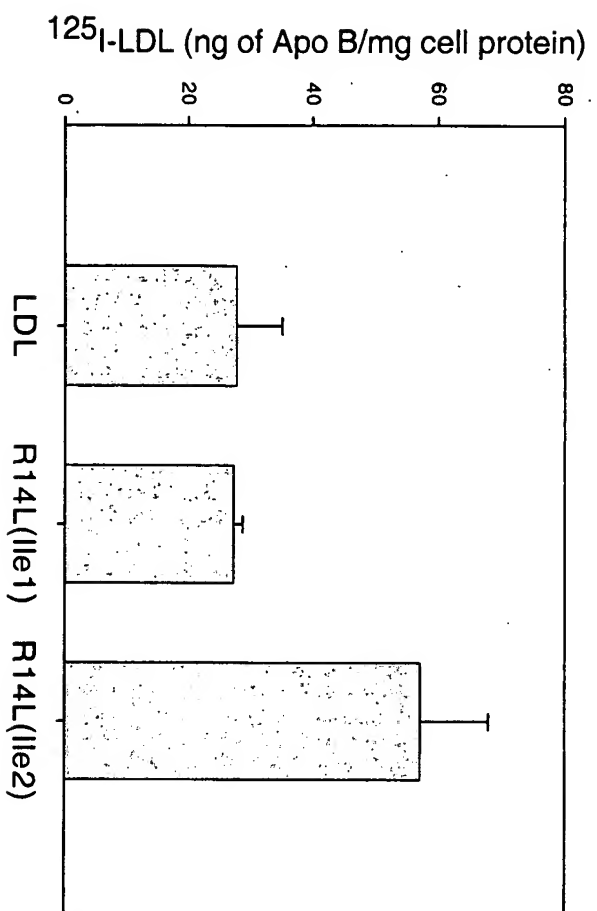


Fig. 9

**Apo E null; i.v. injections of two
single domain cationic peptides**

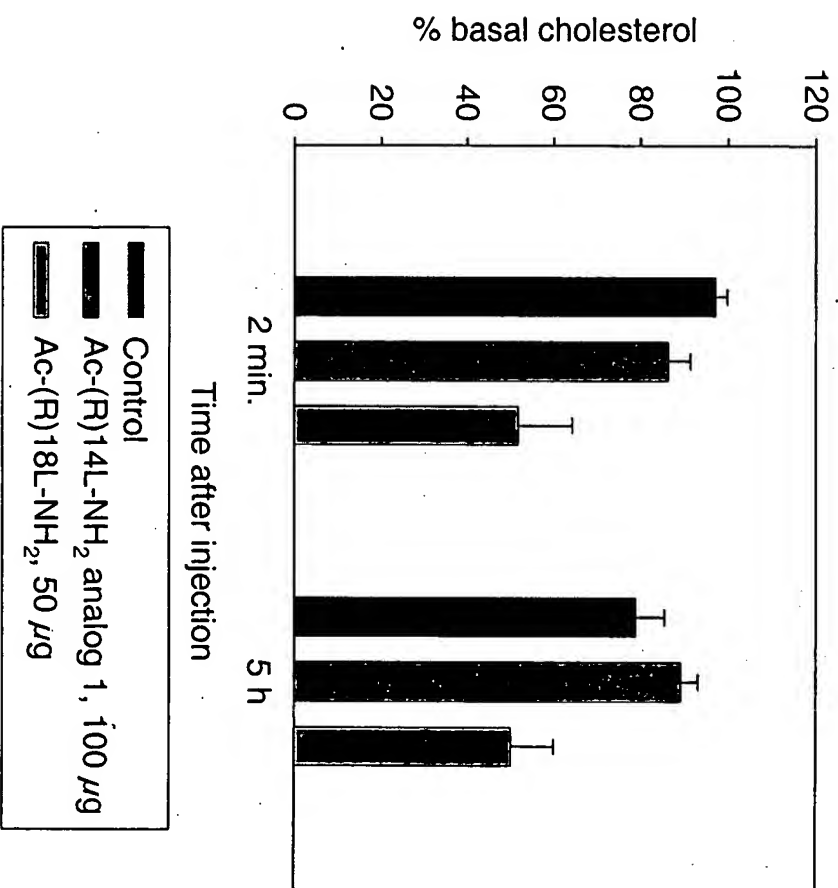


Fig. 10

Apo E null; 100 μ g peptide i.v.
Ac-(R)14L-NH₂ analog 2

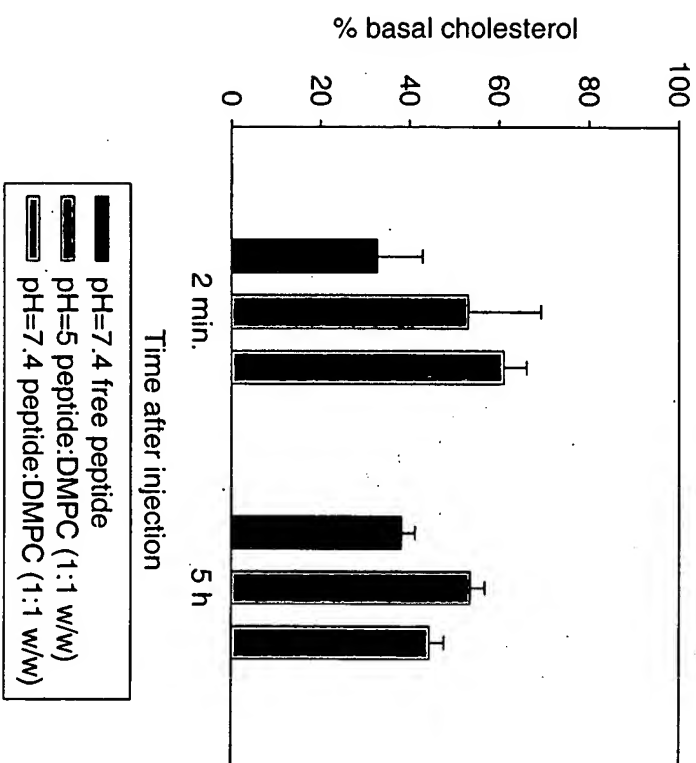
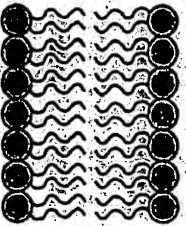
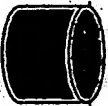
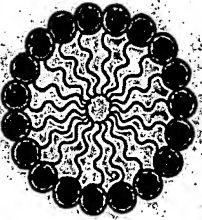

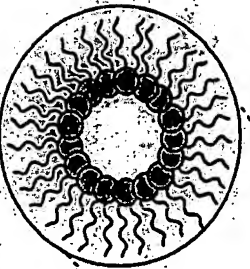



Fig. 11

Figs. 12A-12F

Phase	Molecular shape
 <p>Bilayer</p>	 <p>Cylindric</p>
 <p>Micellar</p>	 <p>Inverted cone</p>
 <p>Hexagonal (H_{II})</p>	 <p>Cone</p>